#73

1600

RAW SEQUENCE LISTING DATE: 02/28/2002
PATENT APPLICATION: US/09/828,995B TIME: 13:15:50

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02282002\I828995B.raw

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      3 <110> APPLICANT: Heska Corporation
      4       McCall, Catherine A.
      5       Tang, Liang A.
      7 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS RELATED TO CANINE IgG AND CANINE
IL-13 RECEPTORS
      9 <130> FILE REFERENCE: AL-7
     11 <140> CURRENT APPLICATION NUMBER: 09/828,995B
     12 <141> CURRENT FILING DATE: 2001-04-09
     14 <150> PRIOR APPLICATION NUMBER: 60/195,874
     15 <151> PRIOR FILING DATE: 2000-04-07
     17 <150> PRIOR APPLICATION NUMBER: 60/195,659
     18 <151> PRIOR FILING DATE: 2000-04-07
     20 <160> NUMBER OF SEQ ID NOS: 104
     22 <170> SOFTWARE: PatentIn version 3.1
     24 <210> SEQ ID NO: 1
     25 <211> LENGTH: 51
     26 <212> TYPE: DNA
     27 <213> ORGANISM: Canis familiaris
     29 <220> FEATURE:
     30 <221> NAME/KEY: CDS
     31 <222> LOCATION: (1)..(51)
     32 <223> OTHER INFORMATION:
     35 <400> SEQUENCE: 1
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     37 Phe Asn Glu Cys Arg Cys Thr Asp Thr Pro Pro Cys Pro Val Pro Glu
     38 1               5                   10                  15
     40 cct                                                               51
     41 Pro
     45 <210> SEQ ID NO: 2
     46 <211> LENGTH: 17
     47 <212> TYPE: PRT
     48 <213> ORGANISM: Canis familiaris
     50 <400> SEQUENCE: 2
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     53 1               5                   10                  15
     56 Pro
     60 <210> SEQ ID NO: 3
     61 <211> LENGTH: 51
     62 <212> TYPE: DNA
     63 <213> ORGANISM: Canis familiaris
     65 <400> SEQUENCE: 3
     66 aggttcaggg actgggcatg ggggtgtatc agtgcatctg cattcattga a            51
     69 <210> SEQ ID NO: 4
     70 <211> LENGTH: 1654
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72 <213> ORGANISM: Canis familiaris
74 <220> FEATURE:
75 <221> NAME/KEY: CDS
76 <222> LOCATION: (70)..(1473)
77 <223> OTHER INFORMATION:
80 <400> SEQUENCE: 4
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83 gacaccacc atg gag tct gtg ttc tgc tgg gtt ttc ctt gtc gtt att tta    111
84           Met Glu Ser Val Phe Cys Trp Val Phe Leu Val Val Ile Leu
85           1               5                  10
87 aaa ggt gtc cag ggt gag gtg cag ttg gtg gag tct ggg gga gac ctg     159
88 Lys Gly Val Gln Gly Glu Val Gln Leu Val Glu Ser Gly Gly Asp Leu
89 15                  20                  25                  30
91 gtg aag cct ggg ggg tcc ctg aga ctc tcc tgt gtg gcc tct gga ttc     207
92 Val Lys Pro Gly Gly Ser Leu Arg Leu Ser Cys Val Ala Ser Gly Phe
93                 35                  40                  45
95 acc ttc agt tcg tac tac atg cat tgg atc cgc cag gct cca ggg aag     255
96 Thr Phe Ser Ser Tyr Tyr Met His Trp Ile Arg Gln Ala Pro Gly Lys
97             50                  55                  60
99 ggg ctt cag cgg gtc gca cat att aga ggt gat gga agg act aca cac     303
100 Gly Leu Gln Arg Val Ala His Ile Arg Gly Asp Gly Arg Thr Thr His
101         65                  70                  75
103 tac gca gac gct atg aag ggc cga ttc acc atc tcc aga gac aac gcc     351
104 Tyr Ala Asp Ala Met Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala
105     80                  85                  90
107 aag aac acg ctg tat ctg cag atg aat agc ctg aca gtc gaa gac acg     399
108 Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Thr Val Glu Asp Thr
109 95                  100                 105                 110
111 gct att tat tac tgt gta aag gac ata tac tat ggg gtc ggg gac tat     447
112 Ala Ile Tyr Tyr Cys Val Lys Asp Ile Tyr Tyr Gly Val Gly Asp Tyr
113                 115                 120                 125
115 tgg ggc cag gga acc ctg gtc acc gtc tcc tca gcc tcc acc acg gcc     495
116 Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Thr Ala
117             130                 135                 140
119 ccc tcg gtt ttc cca ctg gcc ccc agc tgc ggg tcc act tcc ggc tcc     543
120 Pro Ser Val Phe Pro Leu Ala Pro Ser Cys Gly Ser Thr Ser Gly Ser
121         145                 150                 155
123 acg gtg gcc ctg gcc tgc ctg gtg tca ggc tac ttc ccc gag cct gta     591
124 Thr Val Ala Leu Ala Cys Leu Val Ser Gly Tyr Phe Pro Glu Pro Val
125     160                 165                 170
127 act gtg tcc tgg aat tcc ggc tcc ttg acc agc ggt gtg cac acc ttc     639
128 Thr Val Ser Trp Asn Ser Gly Ser Leu Thr Ser Gly Val His Thr Phe
129 175                 180                 185                 190
131 ccg tcc gtc ctg cag tcc tca ggg ctt cac tcc ctc agc agc atg gtg     687
132 Pro Ser Val Leu Gln Ser Ser Gly Leu His Ser Leu Ser Ser Met Val
133                 195                 200                 205
135 aca gtg ccc tcc agc agg tgg ccc agc gag acc ttc acc tgc aac gtg     735
136 Thr Val Pro Ser Ser Arg Trp Pro Ser Glu Thr Phe Thr Cys Asn Val
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139 gtc cac cca gcc agc aac act aaa gta gac aag cca gtg ttc aat gaa      783
140 Val His Pro Ala Ser Asn Thr Lys Val Asp Lys Pro Val Phe Asn Glu
141         225                 230                 235
143 tgc aga tgc act gat aca ccc cca tgc cca gtc cct gaa cct ctg gga      831
144 Cys Arg Cys Thr Asp Thr Pro Pro Cys Pro Val Pro Glu Pro Leu Gly
145     240                 245                 250
147 ggg cct tcg gtc ctc atc ttt ccc ccg aaa ccc aag gac atc ctc agg      879
148 Gly Pro Ser Val Leu Ile Phe Pro Pro Lys Pro Lys Asp Ile Leu Arg
149 255                 260                 265                 270
151 att acc cga aca ccc gag gtc acc tgt gtg gtg tta gat ctg ggc cgt      927
152 Ile Thr Arg Thr Pro Glu Val Thr Cys Val Val Leu Asp Leu Gly Arg
153                 275                 280                 285
155 gag gac cct gag gtg cag atc agc tgg ttc gtg gat ggt aag gag gtg      975
156 Glu Asp Pro Glu Val Gln Ile Ser Trp Phe Val Asp Gly Lys Glu Val
157             290                 295                 300
159 cac aca gcc aag acc cag tct cgt gag cag cag ttc aac ggc acc tac     1023
160 His Thr Ala Lys Thr Gln Ser Arg Glu Gln Gln Phe Asn Gly Thr Tyr
161         305                 310                 315
163 cgt gtg gtc agc gtc ctc ccc att gag cac cag gac tgg ctc aca ggg     1071
164 Arg Val Val Ser Val Leu Pro Ile Glu His Gln Asp Trp Leu Thr Gly
165     320                 325                 330
167 aag gag ttc aag tgc aga gtc aac cac ata gac ctc ccg tct ccc atc     1119
168 Lys Glu Phe Lys Cys Arg Val Asn His Ile Asp Leu Pro Ser Pro Ile
169 335                 340                 345                 350
171 gag agg acc atc tct aag gcc aga ggg agg gcc cat aag ccc agt gtg     1167
172 Glu Arg Thr Ile Ser Lys Ala Arg Gly Arg Ala His Lys Pro Ser Val
173                 355                 360                 365
175 tat gtc ctg ccg cca tcc cca aag gag ttg tca tcc agt gac aca gtc     1215
176 Tyr Val Leu Pro Pro Ser Pro Lys Glu Leu Ser Ser Ser Asp Thr Val
177             370                 375                 380
179 agc atc acc tgc ctg ata aaa gac ttc tac cca cct gac att gat gtg     1263
180 Ser Ile Thr Cys Leu Ile Lys Asp Phe Tyr Pro Pro Asp Ile Asp Val
181         385                 390                 395
183 gag tgg cag agc aat gga cag cag gag ccc gag agg aag cac cgc atg     1311
184 Glu Trp Gln Ser Asn Gly Gln Gln Glu Pro Glu Arg Lys His Arg Met
185     400                 405                 410
187 acc ccg ccc cag ctg gac gag gac ggg tcc tac ttc ctg tac agc aag     1359
188 Thr Pro Pro Gln Leu Asp Glu Asp Gly Ser Tyr Phe Leu Tyr Ser Lys
189 415                 420                 425                 430
191 ctc tct gtg gac aag agc cgc tgg cag cag gga gac ccc ttc aca tgt     1407
192 Leu Ser Val Asp Lys Ser Arg Trp Gln Gln Gly Asp Pro Phe Thr Cys
193                 435                 440                 445
195 gcg gtg atg cat gaa act cta cag aac cac tac aca gat cta tcc ctc     1455
196 Ala Val Met His Glu Thr Leu Gln Asn His Tyr Thr Asp Leu Ser Leu
197             450                 455                 460
199 tcc cat tct ccg ggt aaa tgagcaacac gcccggcacc cagcaagccc           1503
200 Ser His Ser Pro Gly Lys
201         465
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PATENT APPLICATION: US/09/828,995B TIME: 13:15:50

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02282002\I828995B.raw

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cccacccttg gctctcagga tcccctgagg acacctgagc ccctgtccct gtgtacataa   1563
ccctgggtag gcacccatca tgaaataaag cacccagcac tgccctgggc cctgcaaaaa   1623
aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa a                                  1654
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<211> LENGTH: 468
<212> TYPE: PRT
<213> ORGANISM: Canis familiaris
<400> SEQUENCE: 5
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Val Gln Gly Glu Val Gln Leu Val Glu Ser Gly Gly Asp Leu Val Lys
            20                  25                  30
Pro Gly Gly Ser Leu Arg Leu Ser Cys Val Ala Ser Gly Phe Thr Phe
        35                  40                  45
Ser Ser Tyr Tyr Met His Trp Ile Arg Gln Ala Pro Gly Lys Gly Leu
    50                  55                  60
Gln Arg Val Ala His Ile Arg Gly Asp Gly Arg Thr Thr His Tyr Ala
65                  70                  75                  80
Asp Ala Met Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn
                85                  90                  95
Thr Leu Tyr Leu Gln Met Asn Ser Leu Thr Val Glu Asp Thr Ala Ile
            100                 105                 110
Tyr Tyr Cys Val Lys Asp Ile Tyr Tyr Gly Val Gly Asp Tyr Trp Gly
        115                 120                 125
Gln Gly Thr Leu Val Thr Val Ser Ser Ala Ser Thr Thr Ala Pro Ser
    130                 135                 140
Val Phe Pro Leu Ala Pro Ser Cys Gly Ser Thr Ser Gly Ser Thr Val
145                 150                 155                 160
Ala Leu Ala Cys Leu Val Ser Gly Tyr Phe Pro Glu Pro Val Thr Val
                165                 170                 175
Ser Trp Asn Ser Gly Ser Leu Thr Ser Gly Val His Thr Phe Pro Ser
            180                 185                 190
Val Leu Gln Ser Ser Gly Leu His Ser Leu Ser Ser Met Val Thr Val
        195                 200                 205
Pro Ser Ser Arg Trp Pro Ser Glu Thr Phe Thr Cys Asn Val Val His
    210                 215                 220
Pro Ala Ser Asn Thr Lys Val Asp Lys Pro Val Phe Asn Glu Cys Arg
225                 230                 235                 240
Cys Thr Asp Thr Pro Pro Cys Pro Val Pro Glu Pro Leu Gly Gly Pro
                245                 250                 255
Ser Val Leu Ile Phe Pro Pro Lys Pro Lys Asp Ile Leu Arg Ile Thr
            260                 265                 270
Arg Thr Pro Glu Val Thr Cys Val Val Leu Asp Leu Gly Arg Glu Asp
        275                 280                 285
Pro Glu Val Gln Ile Ser Trp Phe Val Asp Gly Lys Glu Val His Thr
    290                 295                 300
Ala Lys Thr Gln Ser Arg Glu Gln Gln Phe Asn Gly Thr Tyr Arg Val
305                 310                 315                 320
Val Ser Val Leu Pro Ile Glu His Gln Asp Trp Leu Thr Gly Lys Glu
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PATENT APPLICATION: US/09/828,995B TIME: 13:15:50

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02282002\I828995B.raw

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298                 325                 330                 335
301 Phe Lys Cys Arg Val Asn His Ile Asp Leu Pro Ser Pro Ile Glu Arg
302             340                 345                 350
305 Thr Ile Ser Lys Ala Arg Gly Arg Ala His Lys Pro Ser Val Tyr Val
306         355                 360                 365
309 Leu Pro Pro Ser Pro Lys Glu Leu Ser Ser Ser Asp Thr Val Ser Ile
310     370                 375                 380
313 Thr Cys Leu Ile Lys Asp Phe Tyr Pro Pro Asp Ile Asp Val Glu Trp
314 385                 390                 395                 400
317 Gln Ser Asn Gly Gln Gln Glu Pro Glu Arg Lys His Arg Met Thr Pro
318                 405                 410                 415
321 Pro Gln Leu Asp Glu Asp Gly Ser Tyr Phe Leu Tyr Ser Lys Leu Ser
322             420                 425                 430
325 Val Asp Lys Ser Arg Trp Gln Gln Gly Asp Pro Phe Thr Cys Ala Val
326         435                 440                 445
329 Met His Glu Thr Leu Gln Asn His Tyr Thr Asp Leu Ser Leu Ser His
330     450                 455                 460
333 Ser Pro Gly Lys
334 465
337 <210> SEQ ID NO: 6
338 <211> LENGTH: 1654
339 <212> TYPE: DNA
340 <213> ORGANISM: Canis familiaris
342 <400> SEQUENCE: 6
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347 tcctcagggg atcctgagag ccaagggtgg ggggcttgct gggtgccggg cgtgttgctc    180
349 atttacccgg agaatgggag agggatagat ctgtgtagtg gttctgtaga gtttcatgca    240
351 tcaccgcaca tgtgaagggg tctccctgct gccagcggct cttgtccaca gagagcttgc    300
353 tgtacaggaa gtaggacccg tcctcgtcca gctggggcgg ggtcatgcgg tgcttcctct    360
355 cgggctcctg ctgtccattg ctctgccact ccacatcaat gtcaggtggg tagaagtctt    420
357 ttatcaggca ggtgatgctg actgtgtcac tggatgacaa ctcctttggg gatggcggca    480
359 ggacatacac actgggctta tgggccctcc ctctggcctt agagatggtc ctctcgatgg    540
361 gagacgggag gtctatgtgg ttgactctgc acttgaactc cttccctgtg agccagtcct    600
363 ggtgctcaat ggggaggacg ctgaccacac ggtaggtgcc gttgaactgc tgctcacgag    660
365 actgggtctt ggctgtgtgc acctccttac catccacgaa ccagctgatc tgcacctcag    720
367 ggtcctcacg gcccagatct aacaccacac aggtgacctc gggtgttcgg gtaatcctga    780
369 ggatgtcctt gggtttcggg ggaaagatga ggaccgaagg ccctcccaga ggttcaggga    840
371 ctgggcatgg gggtgtatca gtgcatctgc attcattgaa cactggcttg tctactttag    900
373 tgttgctggc tgggtggacc acgttgcagg tgaaggtctc gctgggccac ctgctggagg    960
375 gcactgtcac catgctgctg agggagtgaa gccctgagga ctgcaggacg gacgggaagg   1020
377 tgtgcacacc gctggtcaag gagccggaat tccaggacac agttacaggc tcggggaagt   1080
379 agcctgacac caggcaggcc agggccaccg tggagccgga agtggacccg cagctggggg   1140
381 ccagtgggaa aaccgagggg gccgtggtgg aggctgagga gacggtgacc agggttccct   1200
383 ggccccaata gtccccgacc ccatagtata tgtcctttac acagtaataa atagccgtgt   1260
385 cttcgactgt caggctattc atctgcagat acagcgtgtt cttggcgttg tctctggaga   1320
387 tggtgaatcg gcccttcata gcgtctgcgt agtgtgtagt ccttccatca cctctaatat   1380
389 gtgcgacccg ctgaagcccc ttccctggag cctggcggat ccaatgcatg tagtacgaac   1440
391 tgaaggtgaa tccagaggcc acacaggaga gtctcaggga ccccccaggc ttcaccaggt   1500
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Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY DATE: 02/28/2002
PATENT APPLICATION: US/09/828,995B TIME: 13:15:51

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02282002\I828995B.raw

L:829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:1139 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1597 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1686 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:1736 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1827 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1829 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1907 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:1909 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30
L:2726 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:2839 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:3458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57
L:3544 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3699 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
L:3823 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62
L:5961 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83
L:5997 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84
L:6027 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85
L:6057 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86
L:6093 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:87
L:6117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:88